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==============================================================
Sequence Listing could not be accepted due to errors.
See attached Validation Report.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: Anne Corrigan
Timestamp: [year=2008; month=8; day=27; hr=16; min=56; sec=30; ms=742; ]
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Reviewer Comments:
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Trp Leu Asp Phe
1




1

Please remove the "1" above, which appears at the end of the submitted
file.

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Application No: 10587529   Version No: 1.0

Input Set:

Output Set:

Started: 2008-08-23 06:05:24.521
Finished: 2008-08-23 06:05:27.929
Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 408 ms
Total Warnings: 25
Total Errors: 6
No. of SeqIDs Defined: 27
Actual SeqID Count: 27

Error code   Error Description

W 213   Artificial or Unknown found in <213> in SEQ ID (1)

W 213   Artificial or Unknown found in <213> in SEQ ID (2)

W 213   Artificial or Unknown found in <213> in SEQ ID (3)

E 257   Invalid sequence data feature in <221> in SEQ ID (3)

W 213   Artificial or Unknown found in <213> in SEQ ID (4)

W 213   Artificial or Unknown found in <213> in SEQ ID (5)

W 213   Artificial or Unknown found in <213> in SEQ ID (6)

E 257   Invalid sequence data feature in <221> in SEQ ID (6)

W 213   Artificial or Unknown found in <213> in SEQ ID (9)

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W 213   Artificial or Unknown found in <213> in SEQ ID (13)

W 213   Artificial or Unknown found in <213> in SEQ ID (14)

W 213   Artificial or Unknown found in <213> in SEQ ID (15)

W 213   Artificial or Unknown found in <213> in SEQ ID (16)

W 213   Artificial or Unknown found in <213> in SEQ ID (17)

W 213   Artificial or Unknown found in <213> in SEQ ID (18)

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              Input Set:


              Output Set:



             Started:   2008-08-23 06:05:24.521
            Finished:   2008-08-23 06:05:27.929
             Elapsed:   0 hr(s) 0 min(s) 3 sec(s) 408 ms
      Total Warnings:   25
        Total Errors:   6
  No. of SeqIDs Defined:   27
  Actual SeqID Count:   27
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Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (21)
W 213	Artificial or Unknown found in <213> in SEQ ID (22) This error has occured more than 20 times, will not be displayed
E 257	Invalid sequence data feature in <221> in SEQ ID (26)
E 257	Invalid sequence data feature in <221> in SEQ ID (27)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (27)

SEQUENCE LISTING

<110>   WARATAH PHARMACEUTICALS, INC.
        CRUZ, Antonio

<120>   METHODS AND COMPOSITIONS USING CD3 AGONISTS

<130>   24492-020 CIP NATL

<140>   10587529
<141>   2008-08-23

<150>   60/584,635
<151>   2004-07-01

<160>   27

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Ser Ser Tyr Leu Glu Gly Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu
            20   25   30

Val Lys Gly Arg Gly
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<220>
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            20   25   30

Val Lys Gly Arg
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<223>   AMIDATION

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            20   25   30

Val Lys Gly Arg
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1   5   10   15

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Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg
            20   25   30

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            20   25   30

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<213> Heloderma horridum

<300>
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<309> 2004-06-15
<313> (1)..(39)

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Ser Gly Ala Pro Pro Pro Ser
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            20   25   30

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Asp Leu Ser Lys Gln Met Glu Glu Glu Ala Val Arg Leu Phe Ile Glu

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1                5                     10                     15

Trp Leu Lys Asn Gly Gly Pro Ser Ser Gly Ala Pro Pro Ser
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<220>
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<222>   (1)..(1)
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<400>   11

Xaa Leu Gly Pro Gln Gly Pro Pro His Leu Val Ala Asp Pro Ser Lys
1                5                     10                     15


Lys Gln Gly Pro Trp Leu Glu Glu Glu Glu Ala Tyr Gly Trp Met
            20                  25              30


Asp Phe


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<220>
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Lys Gln Gly Pro Trp Leu Glu Glu Glu Glu Ala Tyr Gly Trp Leu
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Asp Phe

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Phe

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Phe

<210> 15
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Met Gln Arg Leu Cys Val Tyr Val Leu Ile Phe Ala Leu Ala Leu Ala
1   5   10   15

Ala Phe Ser Glu Ala Ser Trp Lys Pro Arg Ser Gln Gln Pro Asp Ala
            20   25   30

Pro Leu Gly Thr Gly Ala Asn Arg Asp Leu Glu Leu Pro Trp Leu Glu
            35   40   45

Gln Gln Gly Pro Ala Ser His His Arg Arg Gln Leu Gly Pro Gln Gly
        50   55   60

Pro Pro His Leu Val Ala Asp Pro Ser Lys Lys Gln Gly Pro Trp Leu
65   70   75   80

Glu Glu Glu Glu Glu Ala Tyr Gly Trp Met Asp Phe Gly Arg Arg Ser
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Ala Glu Asp Glu Asn
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<210>   16
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<212>   PRT
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<220>
<223>   Synthesized Peptide

<400>   16

Asp Leu Glu Leu Pro Trp Leu Glu Gln Gln Gly Pro Ala Ser His His
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Arg Arg Gln Leu Gly Pro Gln Gly Pro Pro His Leu Val Ala Asp Pro
            20   25   30

Ser Lys Lys Gln Gly Pro Trp Leu Glu Glu Glu Glu Glu Ala Tyr Gly
            35   40   45

Trp Met Asp Phe
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<210> 17
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<220>
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<400> 17

Trp Leu Glu Glu Glu Glu Glu Ala Tyr Gly Trp Met Asp Phe
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<210> 18
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<400> 18

Tyr Gly Trp Met Asp Phe
1   5

<210> 19
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<212> PRT
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<220>
<223> Synthesized Peptide

<400> 19

Tyr Gly Trp Leu Asp Phe
1   5

<210> 20
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            20   25   30

Ser Ser Gly Ala Pro Pro Pro Ser
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<210> 22
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Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
            20   25   30

Ser Gly Ala Pro Pro Ser Lys Lys Lys Lys Lys Lys
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<210>   23
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<220>
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<210>   24
<211>   6
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